

41939

November 8, 2002

Ms. Joan Kessner Bechtel Hanford Inc. 3350 George Washington Way Richland, WA 99352 MSIN: H0-25

Reference:

P.O. #630

Eberline Services R2-10-090-7376, SDG H1939

Dear Ms. Kessner:

Enclosed is the data report for three other solid samples designated under SAF No. B00-055 received at Eberline Services on October 22 and 25, 2002. The samples were analyzed according to the accompanying chain-of-custody documents.

Please call if you have any questions concerning this report.

Sincerely,

Melissa C. Mannion

Allen Mann

Program Manager

MCM

Enclosure: Data Package





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Case Narrative

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1.0 GENERAL

Bechtel Hanford Inc. (BHI) Sample Delivery Group H1939 was composed of three other solid samples designated under SAF No. B00-055 with a Project Designation of: 100-NR-1 TSD Sites R.A. Sampling – Other Solid. The data is in units of picoCurie per sample (pCi/smpl.)

The samples were received as stated on the Chain-of-Custody documents. Any discrepancies are noted on the Eberline Services Sample Receipt Checklist. The results were transmitted to BHI via e-Fax on November 8, 2002.

2.0 ANALYSIS NOTES

2.1 Total Strontium Analyses

No problems were encountered during the course of the analyses.

2.2 Isotopic Plutonium Analyses

No problems were encountered during the course of the analyses.

2.3 Americium-241 Analyses

No problems were encountered during the course of the analyses.

Case Narrative Certification Statement

"I certify that this data package is in compliance with the SOW, both technically and for completeness, for other than the conditions detailed above. Release of the data obtained in this hard copy data package has been authorized by the Laboratory Manager or a designee, as verified by the following signature."

Mulissa C. Mannion
Program Manager

11 | 8/0 z______

E B E R L I N E S E R V I C E S / R I C H M O N D SAMPLE DELIVERY GROUP H1939

SDG 7376
Contact Melissa C. Mannion

Client <u>Hanford</u>
Contract <u>No. 630</u>
Case no <u>SDG H1939</u>

SUMMARY DATA SECTION

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Mulin	Mann	
Prepared by		
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Reviewed by		

SAMPLE DELIVERY GROUP H1939

SDG 7376
Contact Melissa C. Mannion

REPORT GUIDE

Client <u>Hanford</u>
Contract <u>No. 630</u>
Case no <u>SDG H1939</u>

ABOUT THE DATA SUMMARY SECTION

The Data Summary Section of a Data Package has all data, in several useful orders, necessary for first level, routine review of the data package for a Sample Delivery Group (SDG). This section follows the Data Package Narrative, which has an overview of the data package and a discussion of special problems. It is followed by the Raw Data Section, which has full details.

The Data Summary Section has several groups of reports:

SAMPLE SUMMARIES

The Sample and QC Summary Reports show all samples, including QC samples, reported in one SDG. These reports cross-reference client and lab sample identifiers.

PREPARATION BATCH SUMMARY

The Preparation Batch Summary Report shows all preparation batches (lab groupings reflecting how work was organized) relevant to the reported SDG with information necessary to check the completeness and consistency of the SDG.

WORK SUMMARY

The Work Summary Report shows all samples and work done on them relevant to the reported SDG.

METHOD BLANKS

The Method Blank Reports, one for each Method Blank relevant to the SDG, show all results and primary supporting information for the blanks.

LAB CONTROL SAMPLES

The Lab Control Sample Reports, one for each Lab Control Sample relevant to the SDG, show all results, recoveries and primary supporting information for these QC samples.

REPORT GUIDES

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SUMMARY DATA SECTION

Page 1

Version Ver 1.0
Form DVD-RG
Version 3.06
Report date 11/08/02

Lab id EBRLNE
Protocol Hanford

SAMPLE DELIVERY GROUP H1939

SDG 7376
Contact Melissa C. Mannion

GUIDE, cont.

Client <u>Hanford</u>
Contract <u>No. 630</u>
Case no <u>SDG H1939</u>

ABOUT THE DATA SUMMARY SECTION

DUPLICATES

The Duplicate Reports, one for each Duplicate and Original sample pair relevant to the SDG, show all results, differences and primary supporting information for these QC samples.

MATRIX SPIKES

The Matrix Spike Reports, one for each Spiked and Original sample pair relevant to the SDG, show all results, recoveries and primary supporting information for these QC samples.

DATA SHEETS

The Data Sheet Reports, one for each client sample in the SDG, show all results and primary supporting information for these samples.

METHOD SUMMARIES

The Method Summary Reports, one for each test used in the SDG, show all results, QC and method performance data for one analyte on one or two pages. (A test is a short code for the method used to do certain work to the client's specification.)

REPORT GUIDES

The Report Guides, one for each of the above groups of reports, have documentation on how to read the associated reports.

REPORT GUIDES
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Page 2

Lab id EBRLNE
Protocol Hanford
Version Ver 1.0
Form DVD-RG
Version 3.06

Report date <u>11/08/02</u>

SAMPLE DELIVERY GROUP H1939

SDG 7376 Contact Melissa C. Mannion

SAMPLE SUMMARY

Client <u>Hanford</u> Contract No. 630 Case no SDG N1939

CLIENT SAMPLE ID	LOCATION	NATRIX LEVEL	LAB SAMPLE ID	SAF NO	CHAIN OF CUSTODY	COLLECTED
J006K0	100N	FILTERS	R210090-01	B00-055	B00-055-12	10/15/02 12:2
J006K1	100N	FILTERS	R210090-02	B00-055	B00-055-12	10/15/02 14:3
J006K2	100N	FILTERS	R210090-03	B00-055	B00-055-13	10/21/02 14:2
Method Blank		FILTERS	R210090-05	B00-055		
Lab Control Sample		FILTERS	R210090-04	B00-055		
Duplicate (R210090-01)	100N	FILTERS	R210090-06	B00-055		10/15/02 12:2

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SDG 7376
Contact Melissa C. Mannion

QC SUMMARY

Client Hanford

Contract No. 630

Case no SDG H1939

QC BATCH	CHAIN OF CUSTODY	CLIENT SAMPLE ID	MATRIX	% SOLIDS	SAMPLE AMOUNT	BASIS	DAYS SIN	CE LAB	DEPARTMENT SAMPLE ID
7376	B00-055-12	J006K0 J006K1	FILTERS FILTERS		0.17 g 0.18 g			7 R210090-01 7 R210090-02	7376-001 7376-002
	B00-055-13	J006K2	FILTERS	100.0	0.19 g		10/25/02	4 R210090-03	7376-003
		Method Blank Lab Control Sample Duplicate (R210090-01)	FILTERS FILTERS FILTERS	100.0	0.17 g		10/22/02	R210090-05 R210090-04 7 R210090-06	7376-005 7376-004 7376-006

QC SUMMARY
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SAMPLE DELIVERY GROUP H1939

SDG <u>7376</u> Contact <u>Melissa C. Mannion</u>

PREP BATCH SUMMARY

Client	<u>Hanford</u>
Contract	No. 630
Case no	SDG H1939

TEST	MATRIX	METHOD	PREPARATION BATCH		CLIENT		NCHETS .		DUP/ORIG MS/ORIG	QUALI- FIERS
Alpha AM	Spectros FILTERS	copy Americium-241 in Filters	7041-190	5.0	3		1	1	1/1	
PU	FILTERS	Plutonium, Isotopic in Filters	7041-190	5.0	3		1	1	1/1	
Beta (Counting FILTERS	Total Strontium in Filters	7041-190	10.0	3		1	1	1/1	

Duplicates and Matrix Spikes are those with original (Client) sample in this Sample Delivery Group.

Blank and LCS planchets are those in the same preparation batch as some Client, Duplicate or Spike sample.

PREP BATCH SUMMARY
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Lab id <u>EBRLNE</u>

Protocol <u>Hanford</u>

Version <u>Ver 1.0</u>

Form <u>DVD-PBS</u>

Version <u>3.06</u>

Report date <u>11/08/02</u>

SDG <u>7376</u> Contact <u>Melissa C. Mannion</u>

WORK SUMMARY

Client <u>Hanford</u>
Contract <u>No. 630</u>
Case no <u>SDG H1939</u>

CLIENT SAMPLE LOCATION CUSTODY	ID SAF No	MATRIX	LAB SAMPLE III COLLECTED RECEIVED	PLANCHET	TEST	SUF- FIX	ANALYZED	REVIEWED	BY	METHOD
CUSTODY	SAF NO		RECEIVED	PLANUICI	1591	LIV	ARALIZED	KEATEMEN	D1	HE I HOD
J006K0			R210090-01	7376-001	AM		11/06/02	11/08/02	MCM	Americium-241 in Filters
100N		FILTERS	10/15/02	7376-001	PU		11/08/02	11/08/02	MCM	Plutonium, Isotopic in Filters
B00-055-12	B00-055		10/22/02	7376-001	SR		11/07/02	11/08/02	MCM	Total Strontium in Filters
J006K1			R210090-02	7376-002	AM		11/06/02	11/08/02	MCM	Americium-241 in Filters
100N		FILTERS	10/15/02	7376-002	PU		11/08/02	11/08/02	MCM	Plutonium, Isotopic in Filters
800-055-12	800-055		10/22/02	7376-002	SR		11/07/02	11/08/02	MCM	Total Strontium in Filters
J006K2			R210090-03	7376-003	AM		11/06/02	11/08/02	мсм	Americium-241 in Filters
100N		FILTERS	10/21/02	7376-003	PU		11/08/02	11/08/02	MCM	Plutonium, Isotopic in Filters
B00-055-13	800-055		10/25/02	7376-003	SR		11/07/02	11/08/02	MCM	Total Strontium in Filters
Method Blank			R210090-05	7376-005	АМ		11/06/02	11/08/02	MCM	Americium-241 in Filters
		FILTERS		7376-005	PU		11/08/02	11/08/02	MCM	Plutonium, Isotopic in Filters
	в00-055			7376-005	SR		11/07/02	11/08/02	MCM	Total Strontium in Filters
Lab Control Sa	mple		R210090-04	7376-004	AM		11/06/02	11/08/02	MCM	Americium-241 in Filters
		FILTERS		7376-004	PU		11/08/02	11/08/02	MCM	Plutonium, Isotopic in Filters
	в00-055			7376-004	SR		11/07/02	11/08/02	MCM	Total Strontium in Filters
Duplicate (R21	0090-01)		R210090-06	7376-006	AM		11/06/02	11/08/02	MCM	Americium-241 in Filters
100N		FILTERS	10/15/02	7376-006	PU		11/08/02	11/08/02	MCM	Plutonium, Isotopic in Filters
	800-055		10/22/02	7376-006	SR		11/07/02	11/08/02	MCM	Total Strontium in Filters

TEST	SAF No	COUNTS OF	TESTS BY SAM REFERENCE		RE BLANK	LCS	DUP SPIKE	TOTAL
AM	800-055	Americium-241 in Filters	AMCMISO_IE_PLATE_AEA	3	1	1	1	6
PÚ	800-055	Plutonium, Isotopic in Filters	PUISO_PLATE_AEA	3	1	1	1	6
SR	B00-055	Total Strontium in Filters	SRTOT_SEP_PRECIP_GPC	3	1	1	1	6
TOTALS				9	3	3	3	18

WORK SUMMARY
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R210090-05

METHOD BLANK

Method Blank

	7376 Melissa C. Mannion	Client/Case no Contract	Hanford No. 630	SDG H1939
Lab sample id Dept sample id		Client sample id Material/Matrix SAF No		FILTERS

ANALYTE	CAS NO	RESULT pCi/smpl	2σ ERR (COUNT)	MDA pCi/smpl	RDL pCi/smpl	QUALI- FIERS	TEST
Total Strontium	SR-RAD	-0.201	1.5	3.2	1.0	ប	SR
Plutonium 238	13981-16-3	0	0.31	1.2	1.0	ប	PU
Plutonium 239/240	PU-239/240	0	0.31	1.2	1.0	ប	PU
Americium 241	14596-10-2	-0.297	0.20	0.76	1.0	ับ	AM

100-NR-1 TSD Sites R.A. Smpl.-Solid

QC-BLANK 43047

METHOD BLANKS
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SAMPLE DELIVERY GROUP H1939

R210090-04

LAB CONTROL SAMPLE

Lab Control Sample

SDG <u>7376</u>	Client/Case no <u>Hanford</u> <u>SDG H1939</u>
Contact <u>Melissa C. Mannion</u>	Case no <u>No. 630</u>
Lab sample id <u>R210090-04</u> Dept sample id <u>7376-004</u>	Client sample id <u>Lab Control Sample</u> Material/Matrix <u>FILTERS</u> SAF No <u>BOO-055</u>

ANALYTE	RESULT pCi/smpl	2σ ERR (COUNT)	MDA pCi/smpl	RDL pCi/smpl	QUALI- FIERS	TEST	ADDED pCi/smpl	2σ ERR pCi/smpl	REC %	3σ LMTS (TOTAL)	PROTOCOL
Total Strontium	250	11	3,8	1.0		SR	224	9.0	112	81-119	80-120
Plutonium 238	127	17	1.8	1.0		PU	134	5.4	95	79-121	80-120
Plutonium 239/240	143	18	1.8	1.0		PU	145	5.8	99	79-121	80-120
Americium 241	99.6	8.8	0.79	1.0		AM	105	4.2	95	84-116	80-120

100-NR-1 TSD Sites R.A. Smpl.-Solid

QC-LCS 4304	6			

LAB CONTROL SAMPLES
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SAMPLE DELIVERY GROUP H1939

R210090-06

DUPLICATE

J006K0

SDG 7376 Contact Melissa C. Manni		Client/Case no Case no	
DUPLICATE	ORIGINAL Lab sample id R <u>210090-01</u>	Client sample id	100440
Lab sample id <u>R210090-06</u> Dept sample id <u>7376-006</u>	Dept sample id <u>7376-001</u>	Location/Matrix	
% solids 100.0	Received <u>10/22/02</u> % solids <u>100.0</u>	Collected/Weight Custody/SAF No	10/15/02 12:20 0.17 g 800-055-12 800-055

ANALYTE	DUPLICATE pCi/smpl	20 ERR (COUNT)	MDA pCi/smpl	RDL pCi/smpl	QUALI- FIERS	TEST	ORIGINAL pCi/smpl	2σ ERR (COUNT)	MDA pCi/smpl	QUALI- R	PD %	3ø PROT
Total Strontium	38.9	3.6	2.4	1.0		SR	39.6	3.5	2.2		2	29
Plutonium 238	5.79	1.9	1.2	1.0		PÜ	4.75	1.8	1.1		20	75
Plutonium 239/240	35.8	5.4	1.2	1.0		PU	35.1	5.2	1.1		2	33
Americium 241	27.6	3.7	0.79	1.0		AM	28.9	4.5	1.0		5	33

100-NR-1 TSD Sites R.A. Smpl.-Solid

DUPLICATES
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R210090-01

DATA SHEET

J006K0

•	7376 Melissa C. Mannion	Client/Case no Contract		H1939
1		Client sample id Location/Matrix Collected/Weight Custody/SAF No	100N 10/15/02 12:20 0.17 q	FILTERS

ANALYTE	CAS NO	RESULT pCi/smpl	2σ ERR (COUNT)	MDA pCi/smpl	RDL pCi/smpl	QUALI- FIERS	TRST
Total Strontium	SR-RAD	39.6	3.5	2.2	1.0		SR
Plutonium 238	13981-16-3	4.75	1.8	1.1	1.0		PU
Plutonium 239/240	PU-239/240	35.1	5.2	1.1	1.0		ΡŪ
Americium 241	14596-10-2	28.9	4.5	1.0	1.0		AM

100-NR-1 TSD Sites R.A. Smpl.-Solid

DATA SHEETS
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R210090-02

DATA SHEET

J006K1

	7376 Melissa C. Mannion	Client/Case no Contract		H1939
· ·		Client sample id Location/Matrix Collected/Weight Custody/SAF No	100N 10/15/02 14:30 0.18 g	FILTERS

ANALYTE	CAS NO	RESULT pCi/smpl	2σ ERR (COUNT)	MDA pCi/smpl	RDL pCi/smpl	QUALI- FIERS	TRST
Total Strontium	SR-RAD	77.8	3.4	1.1	1.0		SR
Plutonium 238	13981-16-3	8.38	1.7	0.53	1.0		PU
Plutonium 239/240	PU-239/240	60.4	6.0	0.53	1.0		₽U
Americium 241	14596-10-2	64.4	5.6	0.44	1.0		AM

100-NR-1 TSD Sites R.A. Smpl.-Solid

DATA SHEETS
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R210090-03

DATA SHEET

J006K2

•	7376 Melissa C. Mannion	Client/Case no Contract		H1939
1		Client sample id Location/Matrix Collected/Weight Custody/SAF No	100N 10/21/02 14:20 0.19 g	FILTERS

analyte	CAS NO	RESULT pCi/smpl	2σ ERR (COUNT)	MDA pCi/smpl	RDL pCi/smpl	QUALI- FIERS	TEST
Total Strontium	SR-RAD	-0.182	0.60	1.2	1.0	ט	SR
Plutonium 238	13981-16-3	0.558	0.48	0.51	1.0	ប	PU
Plutonium 239/240	PU-239/240	7.41	1.7	0.61	1.0		₽U
Americium 241	14596-10-2	2.04	0.76	0.47	1.0		AM

100-NR-1 TSD Sites R.A. Smpl.-Solid

DATA SHEETS
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SAMPLE DELIVERY GROUP H1939

Test AM Matrix FILTERS SDG 7376 Contact Melissa C. Mannion

METHOD SUMMARY AMERICIUM-241 IN FILTERS ALPHA SPECTROSCOPY

Client <u>Hanford</u> Contract No. 630 Contract SDG H1939

RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW SUF- TEST FIX PLANCHET	Americium 241	
Preparation batch 7041-	190			
J006K0	R210090-01	7376-001	28.9	
J006K1	R210090-02	7376-002	64.4	
J006K2	R210090-03	7376-003	2.04	
BLK (QC ID=43047)	R210090-05	7376-005	υ ·	
LCS (QC 1D=43046)	R210090-04	7376-004	ok	
Duplicate (R210090-01)	R210090-06	7376-006	ok	

100-NR-1 TSD Sites R.A. Smpl.-Solid

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID		SUF- FIX	MDA pCi/sm	• •		DILU-	YIELD			FWHM keV		PREPARED	ANAL- YZED	DETECTO
Preparation batch 7041-1	90 2σ j	orep er	ror 5	.0 %	Reference	Lab	Notebook	7041	pg.	190					
J006K0	R210090-01	t		1.0	0.100			87		128		22	11/05/02	11/06	\$8-011
J006K1	R210090-02	?		0.44	0.200		•	84		156		22	11/05/02	11/06	SS-011
1009KS	R210090-03	i		0.47	0.200			70		157		16	11/05/02	11/06	SS-015
BLK (QC 1D=43047)	R210090-05	5		0.76	0.100			97		158			11/05/02	11/06	SS-043
LCS (QC ID=43046)	R210090-04	•		0.79	0.100			90		157			11/05/02	11/06	S\$-042
Duplicate (R210090-01)	R210090-06	ı		0.79	0.100			86		158		22	11/05/02	11/06	SS-044
(QC ID=43048)												 			
Nominal values and limit	s from meth	od		1.0	0.100			20-105	5	100	100	180			

1	PROCEDURES	REFERENCE	AMCMISO_IE_PLATE_AEA
1		CP-070	Soil Dissolution, < 1.0g Aliquot, rev 5
1		CP-963	Americium and Curium in Water and Dissolved
1			Samples by Extraction Chromatography, rev 3
1		CP-008	Heavy Element Electroplating, rev 7

AVERAGES ± 2 SD MDA 0.71 ± 0.43 FOR 6 SAMPLES YIELD 86 ± 18

METHOD SUMMARIES Page 1 SUMMARY DATA SECTION Page 13

SAMPLE DELIVERY GROUP H1939

Test PU Matrix FILTERS
SDG 7376
Contact Melissa C. Mannion

METHOD SUMMARY PLUTONIUM, ISOTOPIC IN FILTERS ALPHA SPECTROSCOPY

Client <u>Hanford</u>
Contract <u>No. 630</u>
Contract <u>SDG H1939</u>

RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW SUF- TEST FIX PLANCHET	Plutonium 238	Plutonium 239/240	
Preparation batch 7041-	190				
J006K0	R210090-01	7376-001	4.75	35.1	
J006K1	R210090-02	7376-002	8.38	60.4	
J006K2	R210090-03	7376-003	υ	7.41	
BLK (QC ID=43047)	R210090-05	7376-005	, U	U	
LCS (QC ID=43046)	R210090-04	7376-004	ok	ok	
Duplicate (R210090-01)	R210090-06	7376-006	ok	ok	
Nominal values and limit 100-NR-1 TSD Sites R.A.		RDLs (pCi/smpl)	1.0	1.0	

METHOD PERFORMANCE

CLIENT SAMPLE ID	SAMPLE ID		SUF- FIX	MAX MO			DILU- TION	YIELD %	•		FWHM keV	 	PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7041-	190 2ơ p	rep er	ror 5	.0 % R	eference	Lab I	iotebool	7041	pg.	190					
J006K0	R210090~01			1.1	0.100			92		96		24	11/08/02	11/08	SS-067
J006K1	R210090-02			0.53	0.200			92		<u>96</u>		24	11/08/02	11/08	ss-068
J006K2	R210090-03			0.61	0.200			80		<u>95</u>		18	11/08/02	11/08	SS-069
BLK (QC ID=43047)	R210090-05			1,2	0.100			81		<u>96</u>			11/08/01	11/08	ss-072
LCS (QC ID=43046)	R210090-04			1.8	0.100			56		<u>96</u>			11/08/01	11/08	SS-070
Duplicate (R210090-01) (QC ID≈43048)	R210090-06			1.2	0.100			87		<u>95</u>		24	11/08/02	11/08	ss-073
Nominal values and limit	ts from meth	 od		1.0	0.100	 -		20-105	 5	100	100	 180		•	· · ·

	PROCEDURES	REFERENCE	PU1SO_PLATE_AEA
		CP-070	Soil Dissolution, < 1.0g Aliquot, rev 5
	i	CP-941	Plutonium in Water and Dissolved Samples by
			Extraction Chromatography, rev 1
		CP-008	Heavy Element Electropiating, rev 7
ŀ			

AVERAGES ± 2 SD	MDA <u>1.1</u> ± <u>0.93</u>
FOR 6 SAMPLES	YIELD 81 ± 27

METHOD SUMMARIES
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SUMMARY DATA SECTION
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SAMPLE DELIVERY GROUP H1939

Test <u>SR</u> Matrix <u>FILTERS</u> SDG <u>7376</u>

Contact Melissa C. Mannion

METHOD SUMMARY TOTAL STRONTIUM IN FILTERS BETA COUNTING

Client <u>Hanford</u>
Contract <u>No. 630</u>
Contract <u>SDG H1939</u>

RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW SUF- TEST FIX		Total Strontium		
Preparation batch 7041-	190				 	
J009K0	R210090-01		7376-001	39.6		
J006K1	R210090-02		7376-002	77.8		
J006K2	R210090-03		7376-003	U		
BLK (9C ID=43047)	R210090-05		7376-005	ย		
LC\$ (QC ID=43046)	R210090-04		7376-004	ok		
Duplicate (R210090-01)	R210090-06		7376-006	ok		

METHOD	PERF	ORMA	NCE
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CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST		MDA pCi/smpl			DILU-	YIELD					PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7041-	190 2 <i>σ</i> pr	ep eri	ror 1	0.0 % Re	eference	Lab	Notebook	7041	pg.	190					
J006K0	R210090-01			2,2	0.150			87		100		23	11/07/02	11/07	GRB-207
J006K1	R210090-02			1.1	0.300			85		100		23	11/07/02	11/07	GRB-205
J006K2	R210090-03			1.2	0.300			82		100		17	11/07/02	11/07	GRB-208
BLK (QC ID=43047)	R210090-05			3.2	0.100			85		100			11/07/02	11/07	GRB-203
LCS (QC ID=43046)	R210090-04			3.8	0.100			77		100			11/07/02	11/07	GRB-206
Duplicate (R210090-01) (QC ID=43048)	R210090-06			2.4_	0.150			80		100		23	11/07/02	11/07	GRB-204
Nominal values and limit	s from metho	d		1.0	0.100					100	 -	180			

PROCEDURES	REFERENCE	SRTOT_SEP_PRECIP_GPC
Í	CP-070	Soil Dissolution, < 1.0g Aliquot, rev 5
ļ	CP-502	Strontium in Solids, rev 6

AVERAGES ± 2 SD	MDA	2.3	±	_2.1_
FOR 6 SAMPLES	YIELD	83	±	

METHOD SUMMARIES
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SAMPLE DELIVERY GROUP H1939

SDG 7376
Contact Melissa C. Mannion

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Client	Hanford
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SAMPLE SUMMARY

The Sample and QC Summary Reports show all samples, including QC samples, reported in one Sample Delivery Group (SDG).

The Sample Summary Report fully identifies client samples and gives the corresponding lab sample identification. The QC Summary Report shows at the sample level how the lab organized the samples into batches and generated QC samples. The Preparation Batch and Method Summary Reports show this at the analysis level.

The following notes apply to these reports:

- * LAB SAMPLE ID is the lab's primary identification for a sample.
- * DEPARTMENT SAMPLE ID is an alternate lab id, for example one assigned by a radiochemistry department in a lab.
- * CLIENT SAMPLE ID is the client's primary identification for a sample. It includes any sample preparation done by the client that is necessary to identify the sample.
- * QC BATCH is a lab assigned code that groups samples to be processed and QCed together. These samples should have similar matrices.

QC BATCH is not necessarily the same as SDG, which reflects samples received and reported together.

* All Lab Control Samples, Method Blanks, Duplicates and Matrix Spikes are shown that QC any of the samples. Due to possible reanalyses, not all results for all these QC samples may be relevant to the SDG. The Lab Control Sample, Method Blank, Duplicate, Matrix Spike and Method Summary Reports detail these relationships.

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SAMPLE DELIVERY GROUP H1939

SDG 7376
Contact Melissa C. Mannion

REPORT GUIDE

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PREPARATION BATCH SUMMARY

The Preparation Batch Summary Report shows all preparation batches in one Sample Delivery Group (SDG) with information necessary to check the completeness and consistency of the SDG.

The following notes apply to this report:

- * The preparation batches are shown in the same order as the Method Summary Reports are printed.
- * Only analyses of planchets relevant to the SDG are included.
- * Each preparation batch should have at least one Method Blank and LCS in it to validate client sample results.
- * The QUALIFIERS shown are all qualifiers other than U, J, B, L and H that occur on any analysis in the preparation batch. The Method Summary Report has these qualifiers on a per sample basis.

These qualifiers should be reviewed as follows:

- X Some data has been manually entered or modified. Transcription errors are possible.
- P One or more results are 'preliminary'. The data is not ready for final reporting.
- 2 There were two or more results for one analyte on one planchet imported at one time. The results in DVD may not be the same as on the raw data sheets.

Other lab defined qualifiers may occur. In general, these should be addressed in the SDG narrative.

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SAMPLE DELIVERY GROUP H1939

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WORK SUMMARY

The Work Summary Report shows all samples, including QC samples, and all relevant analyses in one Sample Delivery Group (SDG). This report is often useful as supporting documentation for an invoice.

The following notes apply to this report:

- * TEST is a code for the method used to measure associated analytes. Results and related information for each analyte are on the Data Sheet Report. In special cases, a test code used in the summary data section is not the same as in associated raw data. In this case, both codes are shown on the Work Summary.
- * SUFFIX is the lab's code to distinguish multiple analyses (recounts, reworks, reanalyses) of a fraction of the sample. The suffix indicates which result is being reported. An empty suffix normally identifies the first attempt to analyze the sample.
- * The LAB SAMPLE ID, TEST and SUFFIX uniquely identify all supporting data for a result. The Method Summary Report for each TEST has method performance data, such as yield, for each lab sample id and suffix and procedures used in the method.
- * PLANCHET is an alternate lab identifier for work done for one test. It, combined with the TEST and SUFFIX, may be the best link to raw data.
- * For QC samples, only analyses that directly QC some regular sample are shown. The Lab Control Sample, Method Blank, Duplicate, Matrix Spike and Method Summary Reports detail these relationships.
- * The SAS (Special Analytical Services) Number is a client or lab assigned code that reflects special processing for samples, such as rapid turn around. Counts of tests done are lists by SAS number since it is likely to affect prices.

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SAMPLE DELIVERY GROUP H1939

SDG <u>7376</u>
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REPORT GUIDE

Client <u>Hanford</u>
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Case no <u>SDG H1939</u>

DATA SHEET

The Data Sheet Report shows all results and primary supporting information for one client sample or Method Blank. This report corresponds to both the CLP Inorganics and Organics Data Sheet.

The following notes apply to this report:

- * TEST is a code for the method used to measure an analyte. If the TEST is empty, no data is available; the analyte was not analyzed for.
- * The LAB SAMPLE ID and TEST uniquely identify work within the Summary Data Section of a Data Package. The Work Summary and Method Summary Reports further identify raw data that underlies this work.

The Method Summary Report for each TEST has method performance data, such as yield, for each Lab Sample ID and a list of procedures used in the method.

- * ERRORs can be labeled TOTAL or COUNT. TOTAL implies a preparation (non-counting method) error has been added, as square root of sum of squares, to the counting error denoted by COUNT. The preparation errors, which may vary by preparation batch, are shown on the Method Summary Report.
- * A RESULT can be 'N.R.' (Not Reported). This means the lab did this work but chooses not to report it now, possibly because it was reported at another time.
- * When reporting a Method Blank, a RESULT can be 'N.A.' (Not Applicable). This means there is no reported client sample work in the same preparation batch as the Blank's result. This is likely to occur when the Method Blank is associated with reanalyses of selected work for a few samples in the SDG.

The following qualifiers are defined by the DVD system:

U The RESULT is less than the MDA (Minimum Detectable Activity).

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SAMPLE DELIVERY GROUP H1939

SDG <u>7376</u> Contact <u>Melissa C. Mannion</u>

GUIDE, cont.

Client <u>Hanford</u>
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DATA SHEET

If the MDA is blank, the ERROR is used as the limit.

- J The RESULT is less than the RDL (Required Detection Limit) and no U qualifier is assigned.
- B A Method Blank associated with this sample had a result without a U flag and, after correcting for possibly different aliquots, that result is greater than or equal to the MDA for this sample.

Normally, B is not assigned if U is. When method blank subtraction is shown on this report, B flags are assigned based on the unsubtracted values while U's are assigned based on the subtracted ones. Both flags can be assigned in this case.

For each sample result, all Method Blank results in the same preparation batch are compared. The Method Summary Report documents this and other QC relationships.

- L Some Lab Control Sample that QC's this sample had a low recovery. The lab can disable assignment of this qualifier.
- H Similar to 'L' except the recovery was high.
- P The RESULT is 'preliminary'.
- X Some data necessary to compute the RESULT, ERROR or MDA was manually entered or modified.
- 2 There were two or more results available for this analyte. The reported result may not be the same as in the raw data.

Other qualifiers are lab defined. Definitions should be in the SDG narrative.

The following values are underlined to indicate possible problems:

* An MDA is underlined if it is bigger than its RDL.

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SAMPLE DELIVERY GROUP H1939

SDG 7376
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Client <u>Hanford</u>
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DATA SHEET

- * An ERROR is underlined if the 1.645 sigma counting error is bigger than both the MDA and the RESULT, implying that the MDA may not be a good estimate of the 'real' minimum detectable activity.
- * A negative RESULT is underlined if it is less than the negative of its 2 sigma counting ERROR.
- * When reporting a Method Blank, a RESULT is underlined if greater than its MDA. If the MDA is blank, the 2 sigma counting error is used in the comparison.

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REPORT GUIDE

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LAB CONTROL SAMPLE

The Lab Control Sample Report shows all results, recoveries and primary supporting information for one Lab Control Sample.

The following notes apply to this report:

- * All fields in common with the Data Sheet Report have similar usage. Refer to its Report Guide for details.
- * An amount ADDED is the lab's value for the actual amount spiked into this sample with its ERROR an estimate of the error of this amount.

An amount added is underlined if its ratio to the corresponding RDL is outside protocol specified limits.

- * REC (Recovery) is RESULT divided by ADDED expressed as a percent.
- * The first, computed limits for the recovery reflect:
 - 1. The error of RESULT, including that introduced by rounding the result prior to printing.

If the limits are labeled (TOTAL), they include preparation error in the result. If labeled (COUNT), they do not.

- 2. The error of ADDED.
- 3. A lab specified, per analyte bias. The bias changes the center of the computed limits.
- * The second limits are protocol defined upper and lower QC limits for the recovery.
- * The recovery is underlined if it is outside either of these ranges.

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DUPLICATE

The Duplicate Report shows all results, differences and primary supporting information for one Duplicate and associated Original sample.

The following notes apply to this report:

* All fields in common with the Data Sheet Report have similar usage. This applies both to the Duplicate and Original sample data. Refer to the Data Sheet Report Guide for details.

If the Duplicate has data for a TEST and the lab did not do this test to the Original, the Original's RESULTs are underlined.

* The RPD (Relative Percent Difference) is the absolute value of the difference of the RESULTs divided by their average expressed as a percent.

If both RESULTs are less than their MDAs, no RPD is computed and a '-' is printed.

For an analyte, if the lab did work for both samples but has data for only one, the MDA from the sample with data is used as the other's result in the RPD.

* The first, computed limit is the sum, as square root of sum of squares, of the errors of the results divided by the average result as a percent, hence the relative error of the difference rather than the error of the relative difference. The errors include those introduced by rounding the RESULTs prior to printing.

If this limit is labeled TOT, it includes the preparation error in the RESULTs. If labeled CNT, it does not.

This value reported for this limit is at most 999.

- * The second limit for the RPD is the larger of:
 - 1. A fixed percentage specified in the protocol.

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GUIDE, cont.

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DUPLICATE

- 2. A protocol factor (typically 2) times the average MDA as a percent of the average result. This limit applies when the results are close to the MDAs.
- * The RPD is underlined if it is greater than either limit.
- * If specified by the lab, the second limit column is replaced by the Difference Error Ratio (DER), which is the absolute value of the difference of the results divided by the quadratic sum of their one sigma errors, the same errors as used in the first limit.

Except for differences due to rounding, the DER is the same as the RPD divided by the first RPD limit with the limit scaled to 1 sigma.

* The DER is underlined if it is greater than the sigma factor, typically 2 or 3, shown in the header for the first RPD limit.

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MATRIX SPIKE

The Matrix Spike Report shows all results, recoveries and primary supporting information for one Matrix Spike and associated Original sample.

The following notes apply to this report:

* All fields in common with the Data Sheet Report have similar usage. This applies both to the Spiked and Original sample data. Refer to the Data Sheet Report Guide for details.

If the Spike has data for a TEST and the lab did not do this test to the Original, the Original's RESULTs are underlined.

* An amount ADDED is the lab's value for the actual amount spiked into the Spike sample with its ERROR an estimate of the error of this amount.

An amount is underlined if its ratio to the corresponding RDL is outside protocol specified limits.

- * REC (Recovery) is the Spike RESULT minus the Original RESULT divided by ADDED expressed as a percent.
- * The first, computed limits for the recovery reflect:
 - 1. The errors of the two RESULTs, including those introduced by rounding them prior to printing.

If the limits are labeled (TOTAL), they include preparation error in the result. If labeled (COUNT), they do not.

- 2. The error of ADDED.
- 3. A lab specified, per analyte bias. The bias changes the center of the computed limits.
- The second limits are protocol defined upper and lower QC limits

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MATRIX SPIKE

for the recovery.

These limits are left blank if the Original RESULT is more than a protocol defined factor (typically 4) times ADDED. This is a way of accounting for that when the spike is small compared to the amount in the original sample, the recovery is unreliable.

* The recovery is underlined (out of spec) if it is outside either of these ranges.

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SAMPLE DELIVERY GROUP H1939

SDG <u>7376</u> Contact <u>Melissa C. Mannion</u>

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METHOD SUMMARY

The Method Summary Report has two tables. One shows up to five results measured using one method. The other has performance data for the method. There is one report for each TEST, as used on the Data Sheet Report.

The following notes apply to this report:

* Each table is subdivided into sections, one for each preparation batch. A preparation batch is a group of aliquots prepared at roughly the same time in one work area of the lab using the same method.

There should be Lab Control Sample and Method Blank results in each preparation batch since this close correspondence makes the QC meaningful. Depending on lab policy, Duplicates need not occur in each batch since they QC sample dependencies such as matrix effects.

* The RAW TEST column shows the test code used in the raw data to identify a particular analysis if it is different than the test code in the header of the report. This occurs in special cases due to method specific details about how the lab labels work.

The Lab Sample or Planchet ID combined with the (Raw) Test Code and Suffix uniquely identify the raw data for each analysis.

* If a result is less than both its MDA and RDL, it is replaced by just 'U' on this report. If it is greater than or equal to the RDL but less than the MDA, the result is shown with a 'U' flag.

The J and X flags are as on the data sheet.

- * Non-U results for Method Blanks are underlined to indicate possible contamination of other samples in the preparation batch. The Method Blank Report has supporting data.
- * Lab Control Sample and Matrix Spike results are shown as: ok, No data, LOW or HIGH, with the last two underlined. 'No data'

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SAMPLE DELIVERY GROUP H1939

SDG 7376
Contact Melissa C. Mannion

GUIDE, cont.

Client <u>Hanford</u>
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METHOD SUMMARY

means no amount ADDED was specified. 'LOW' and 'HIGH' correspond to when the recovery is underlined on the Lab Control Sample or Matrix Spike Report. See these reports for supporting data.

- * Duplicate sample results are shown as: ok, No data, or OUT, with the last two underlined. 'No data' means there was no original 'sample data found for this duplicate. 'OUT' corresponds to when the RPD is underlined on the Duplicate Report. See this report for supporting data.
- * If the MDA column is labeled 'MAX MDA', there was more than one result measured by the reported method and the MDA shown is the largest MDA. If not all these results have the same RDL, the MAX MDA reflects only those results with RDL equal to the smallest one.

MDAs are underlined if greater than the printed RDL.

- * Aliquots are underlined if less than the nominal value specified for the method.
- * Prepareation factors are underlined if greater than the nominal value specified for the method.
- * Dilution factors are underlined if greater than the nominal value specified for the method.
- * Residues are underlined if outside the range specified for the method. Residues are not printed if yields are.
- * Yields, which may be gravimetric, radiometric or some type of recovery depending on the method, are underlined if outside the range specified for the method.
- * Efficiencies are underlined if outside the range specified for the method. Efficiencies are detector and geometry dependent so this test is only approximate.

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Lab id EBRLNE
Protocol Hanford
Version Ver 1.0

Form <u>DVD-RG</u> Version <u>3.06</u>

Report date <u>11/08/02</u>

SAMPLE DELIVERY GROUP H1939

SDG 7376
Contact Melissa C. Mannion

GUIDE, cont.

Client Hanford
Contract No. 630
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METHOD SUMMARY

- * Count times are underlined if less than the nominal value specified for the method.
- * Resolutions (as FWHM; Full Width at Half Max) are underlined if greater than the method specified limit.
- * Tracer drifts are underlined if their absolute values are greater than the method specified limit. Tracer drifts are not printed if percent moistures are.
- * Days Held are underlined if greater than the holding time specified in the protocol.
- * Analysis dates are underlined if before their planchet's preparation date or, if a limit is specified, too far after it.

For some methods, ratios as percentages and error estimates for them are computed for pairs of results. A ratio column header like '1÷3' means the ratio of the first result column and the third result column.

Ratios are not computed for Lab Control Sample, Method Blank or Matrix Spike results since their matrices are not necessarily similar to client samples'.

The error estimate for a ratio of results from one planchet reflects only counting errors since other errors should be correlated. For a ratio involving different planchets, if QC limits are computed based on total errors, the error for the ratio allows for the preparation errors for the planchets.

The ratio is underlined (out of spec) if the absolute value of its difference from the nominal value is greater than its error estimate. If no nominal value is specified, this test is not done.

For Gross Alpha or Gross Beta results, there may be a column showing the sum of other Alpha or Beta emitters. This sum includes all relevant

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GUIDE, cont.

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METHOD SUMMARY

results in the DVD database, whether reported or not. Results in the sum are weighted by a particles/decay value specified by the lab for each relevant analyte. Results less than their MDA are not included. No sums are computed for Lab Control, Method Blank or Matrix Spike samples since their various planchets may not be physically related.

If a ratio of total isotopic to Gross Alpha or Beta is shown, the error for the ratio reflects both the error in the Gross result and the sum, as square root of sum of squares, of the errors in the isotopic results.

For total elemental uranium or thorium results, there may be a column showing the total weight computed from associated isotopic results. Ignoring results less than their MDAs, this is a weighted sum of the isotopic results. The weights depend on the molecular weight and half-life of each isotope so as to convert activities (decays) to weight (atoms).

If a ratio of total computed to measured elemental uranium or thorium is shown, the error for the ratio reflects the errors in all the measurements.

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Bechtel Hanford Inc.	CI	CHAIN OF CUSTODY/SAMPLE ANALYSIS					REQUEST B0			0-055-1	2 Page	1 0	of <u>1</u>			
B.MILLS C. LANDAS		Company Contact Telephone No. J.D. Fancher 3/3-9123					Project Coordinator TRENT, SJ		OF P	rice Code	7K (₹ ८ ₽	ta Turi	around		
Project Designation 100-NR-1 TSD Sites R. A. Sampling - Other Solid		Sampling Location H/939 (7376							A	ir Qualit	y 🗆	21	Day	THE		
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Potentially Radioactive		Preservation	None				•	f				{		1	1	
Special Handling and/or Storage		Type of Container	Petri Dish													
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FINAL SAMPLE Disposal Method DISPOSITION					Dispos	sed By	<u> </u>						Date/Tit	he		

Bechtel Hanford Inc.			CHAIN OF CUSTODY/SAMPLE ANALYSIS					REQUEST B00-055			0-055-13	55-13 Page <u>1</u> of <u>1</u>	
Collector (7)	Lowery		Company Contact Telephone No. J.D. Fancher 313-9123					Project Coordinator TRENT, SI		Price Code TK GC Data Turuaro			ruaround
Project Designation	tes R. A. Sampling - Other Solid		Sampling Location H1939 (7376)					SAF No. B00-055		Air Quality 18/15 1			day
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Potentially Radioac	LE HAZARDS/REMARKS Hive A Hech A Report		Preservation	None Petri Dish									
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Sample No	o. Matrix •	Sample Date	Sample Time										
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			ed By/Stored In Date/Time			-						L=Liquid V=Vegetation X=Other	
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FINAL SAMPLE DISPOSITION	Disposal Method	·				Dispo	osed By				I	Pate/Time	<u></u>

Richmond, CA Laboratory

SAMPLE RECEIPT CHECKLIST

SAMPLE RECEIPT										
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CoC N	o	B 00	-055 -	12			-T_			
Contai	o	ERL 96	-079 1	Reguested	TAT (Days)	21 Day	.O. Receive	d Yes	[] No []	
				INSP	CTION					
1.	Custody s	eals on shi	pping cont	ajner intact	:7	Yes [🎢	No []	N/A []	
2.	Custody se	eals on shi	oping cont	ainer dated	& signed?	Yes [49	Na [1	N/A[]	
з.	Custody a	eals on san	nple contai	ners intact	?	Yes [Y]	No [}	N/A [1	
4.	Custodys	eals on san	npie contai	ners dated	& signed?	Yes [T]	No []	N/A []	
5.	Packing m	aterial is:				Wet { }	Dry [ya		
6.	Number of	samples ir	shipping	container:	<u> </u>			,		
7.	Number of	containers	per sampl	le:	 	(Or see Co	c)		
8.	Paperwork	agrees wi	th samples	?		Yes [7]	No [}		
9.	Samples h	ave: Tape	[] Heza	rd labels [) Rad lab	els [] App	propriate sa	mple la	ibels [🞾	
10.	Samples a	re: In god	d conditio	n[XP Le	aking []	Broken C	ontainer (] Mis	sing []	
11.	Describe a	ny anomali	es:	 -						
					(K)		<u> </u>			
				ho	<i>//</i>					
13.	Was P.M.	notified of	any anom	elies? Yes			Date			
14.	Received b	γ	Aug.)	mo	Date:	10/22/2	Time:	1010	0 800	
	er Sample No.	cpm .	mr/hr	wipe	Custome No	r Sampie o	cpm	-mr/hr	wipe	
			•							
			·							
									<u> </u>	
										
										
									- 	
<u></u>										
on Chan	nber Ser. No	o			Calibratio	on date	·			
Alpha me	eter Ser. No	·			Calibratio	on date				
Survey N	deter Ser. N	lo		<u></u>	Calibratio	on date		· · ·		